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Q13370; 000639; Q14408;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
20-AUG-2001 (Rel. 46, Last annotation update)
20-AUG-2001 (Rel. 40, Last annotation update)
CGMP_INHIBITED 3',5'-CYCLIC PHOSPHODIESTERASE B (EC 3.1.4.17) (CYCLIC GMP_INHIBITED PHOSPHODIESTERASE B) (CGI-PDE B) (CGIPDE1) (CGIP1).
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"Molecular cloning and chromosomal assignment of the human
of the rat cGMP-inhibited phosphodiesterase 1 (PDE3A) -- a
involved in fat metabolism located at 11p15.1.";
Genomics 37:211-218(1996).
-- FUNCTION: MAY PLAY A ROLE IN FAT METABOLISM.
-- CATALYTIC ACTIVITY: GDRANOSINE 3',5'-CYCLIC PHOSPHATE +
GUANOSINE 5'-PHOSPHATE.
-- ENZYME REGULATION: INHIBITED BY CGMP.
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MIRITINE-97038690; PubMed-8884271;
MIRITINE-97038690; PubMed-8884271;
MIRITINE-97038690; PubMed-8884271;
Ward D., Taira M., Makino H., Manganlello V.C.;
"Characterization of the cDNA and gene encoding human PDE3B, the "CGIPI isoform of the human cyclic GMP-inhibited cyclic nucleotide phosphodiseterase family.";
Genomics 36:476-485(1996).
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Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Pred. No.
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ANIDATION (G-144 PROVIDE AMIDE GROUP).
E3B1CBBE55C09911 CRC64;
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RESULT 3
ADRO_BOVIN
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EMBL; D50632; BAA09306.1; JOINED.
EMBL; D50633; BAA09306.1; JOINED.
EMBL; D50634; BAA09306.1; JOINED.
EMBL; D50634; BAA09306.1; JOINED.
EMBL; D50637; BAA09306.1; JOINED.
EMBL; D50638; BAA09306.1; JOINED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                          01-AUG-1988 (Rel. 08, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation update)
NADPH:ADRENODOXIN OXIDOREDUCTASE, MITOCHONDRIAL PRECURSOR
(EC 1.18.1.2) (ADRENODOXIN REDUCTASE) (AR) (FERREDOXIN-NAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                       MEDIINE=94177140; PubMed=8130767;
Takata Y., Sagara Y., Kono A., Sekimizu K., Hori
"Gene structure of bovine adrenodoxin reductase.
Biol. Pharm. Bull. 16:1200-1206(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                        FDXR OR ADXR
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                                                                of adrenodoxin oxidoreductase from bovine adrenal c Biochem. Biophys. Res. Commun. 145:1239-1247(1987).
                                                                                                                                                                                                                                                                                                                                                                       Bovidae; Bovinae;
           MEDLINE=89170752; PubMed=2924777;
                         SEQUENCE FROM N.A. TISSUE=Adrenal cortex;
                                                                                       Molecular cloning and sequence analysis of full-length cDNA for mRNA
                                                                                                         Yamano T., Okamoto
                                                                                                                      Nonaka Y., Murakami
                                                                                                                                    SEQUENCE FROM N.A.
MEDLINE=87270696; PubMed=3038094;
                                                                                                                                                                                        "Cioning and sequence analysis of adrenodoxin reductase cDNA from bovine adrenal cortex.";
                                                                                                                                                                                                                  Sagara Y., Takata Y., Miyata T.,
                                                                                                                                                                                                                                MEDLINE-88198050;
                                                                                                                                                                                                                                              SEQUENCE FROM N.A.,
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                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9913;
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                                                                                                                                                                           . Biochem. 102:1333-1336(1987).
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5; Conser
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Gutfinger
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                                                                                                                                                                                                                                                AND PARTIAL SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                   Cetartiodactyla;
                                                                                                                                                                                                                                                                                                                                                                                      Chordata; Craniata; Vertebrata; Eute Cetartiodactyla; Ruminantia; Pecora;
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87
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62.5%;
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A -> V (IN REF. 2).
MW; E5141C3DA12E99B0 CRC64;
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Pred. No.
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                                                                                                                       Υ.,
                                                                                                                                                                                                                      Hara T.,
                                                                                                                          Kuramitsu S., Kagamiyama
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AR) (FERREDOXIN-NADP(+)
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EMBL; D00211; BAA00150.1; --
EMBL; X13736; CAA32002.1; --
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by non-profit institutions as long as its content is in no way entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                                                                                                                                                                                                      lectron transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tiegler G.A., Vonrhein C., Hanukoglu I., Schulz G.E.,
"The Structure of adrenodoxin reductase of mitochondrial P450 systems:
electron transfer for steroid biosynthesis.",
J. Mol. Biol. 289; 981-990(1999).
-1- FUNCTION: SERVES AS THE FIRST ELECTRON TRANSFER PROTEIN IN ALL THE MITOCHONDRIAL P450 SYSTEMS: INCLUDING CHOLESTEROL SIDE CHAIN HYDROXYLATION IN THE BARENAL CORTEX, 25-OH-VITAMIN D3-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -1- CATALYTIC ACTIVITY: REDUCED ADRENODOXIN + NADP(+) = OXIDIZED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SUBCELLULAR LOCATION: MITOCHONDRIAL MATRIX.

1- ALTERNATIVE PRODUCTS: 2 ISOFORMS; A SHORT FORM (SHOWN HERE) AND A LONG FORM; ARE PRODUCED BY ALTERNATIVE SPLICING. THE LONG FORM REPRESENTS 10-20% OF ALL ADRENODOXIN REDUCTASE MENA. AND SEEMS TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Adrenal gland;
MEDLINE=99299392; PubMed=10369776;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hanukoglu I., Gutfinger T., Haniu M., Shively J.E.;
"Isolation of a cDNA for adrenodoxin reductase (ferredoxin-NADP+
reductase). Implications for mitochondrial cytochrome P-450 systa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eur. J. Biochem. 169:449-455(1987).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; A29604; A29604.
; JS0390; JS0390.
; S03558; S03558.
; JT0751; JT0751.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HYDROXYLATION IN THE KIDNEY, AND STEROL C-27 HYDROXYLATION IN THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IPR000759; Adrndx_redctse.
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02-JUN-00.
                                                                                                                                                                                                                                                                                                                                                                                     Transit peptide;
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                                                                                                                                                                                                                                                                                                                                                                               Oxidoreductase; Flavoprotein; NADP; FAD;
sit peptide; Alternative splicing; 3D-structure.
                                                                                                            54338 MW;
          52.0%;
                                                                            K -> K (IN REF. 3).

PS -> RL (IN REF. 3).

RAAGIRLAVIR -> ARRSAWOSPE (IN REF. 3).

""" AVIDTGDVEDL -> HPGSAHWCCGGP (IN REF. 3).
Score 51.5; D
Pred. No. 5.4;
                                                                                                                                                                                         QDAYH -> RVYRLT (IN REF. 3).
K -> R (IN REF. 3).
                                                                                                                                                                                                                                                                                                           NADPH:ADRENODOXIN OXIDOREDUCTASE.
E -> EVLLLCQ (IN LONG ISOFORM).
G -> R (IN REF. 3).
                                                                                                                                                                                                                                                          G -> R (IN REF. 3).
FGVAPDHPEVKNVI -> VWLALTTPRSRMLL (IN REF
                        DB 1;
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                                                                                                                                                                                                        CARBOHYD
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                                                                                                                                                                                                                                                                                                                                   EMBL; X16816; CAA34723.1; ..
                                                                                                                                                                                                                                                                                                                                                                       the European Bioinformatics Institute. The by non-profit institutions as long modified and this statement is not removed.
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or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                              InterPro; IPR002551; Corona_S1.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Raabe T., Schelle-Prinz B., Siddell S.G.;

Nucleotide sequence of the gene encoding the spike glycoprotein of human coronavirus HCV 229g.",

J. Gen. Virol. 71:1065-1073(1990).

-I- FUNCTION: THE PEPLOMER PROTEIN MEDIATES THE BINDING OF VIRIONS AND IN SYNCYTIUM FORMATION.

AND IN SYNCYTIUM FORMATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human coronavirus (strain 229E).
Viruses; ssRNA positive-strand viruses, no DNA stage; Nidovirales;
Coronaviridae; Coronavirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P15423;
01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=90264837; PubMed=2345367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E2 GLYCOPROTEIN PRECURSOR (SPIKE GLYCOPROTEIN) (PEPLOMER PROTEIN).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VGL2_CVH22
                                                                                                                                                                                                                                                                                                                            A34766; VGIHHC.
                                                                                                                                                                                                                                                                                                                                                                                                                          SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.
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                N-LINKED
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01-OCT-1996 (Rel. 34, Last sequence update)
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REACTION CENTER PROTEIN L CHAIN (PHOTOSYNTHETIC REACTION CENTER L
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Bacteria; Proteobacteria; beta subdivision; Comamonad
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=28068;
                                                                                                                                                                                                                                                                                                                                                                                                                             Nagashima K.V.P., Matsuura K., Ohyama S., Shimada K.; Primary structure and transcription of genes encoding B870 and
                                                                                                                                                                                                                                     deletion strain.";
Mol. Gen. Genet. 252:379-385(1996).
Mol. Gen. Genet. 252:379-385(1996).
MEDIATES THE INITIAL PHOTOCHEMICAL EVENT IN THE ELECTRON TRANSIMEDIATES THE INITIAL PHOTOCHEMICAL EVENT IN THE ELECTRON TRANSIMEDIATES.
                                                                                                                                                                                                                                                                                                    Ouchane S., Picaud M., Reiss-Husson F., Vernotte C., Astier C.; "Development of gene transfer methods for Rubrivivax gelatinosus S1:
                                                                                                                                                                                                                                                                                                                                                                                                      gelatinosus.";
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                                                                                                                                                                                                                                                                                                                                  MEDLINE=97033541; PubMed=8879238;
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                       the European Bioinformatics Institute. There are no restrictions on use by non-profit institutions as long as its content is in no modified and this statement is not removed. Usage by and for commer entities requires a license agreement (See http://www.isb-sib.ch/announ or send an email to license@isb-sib.ch).
                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
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EMBL; D16822; BAA04100.1; -.
                                                                                                                                                                                                                                                                                                                                                                                         Biol. Chem. 269:2477-2484(1994).
                                                                                                                                                                    TWO BACTERIOPHEOPHYTINS, TWO UBIQUINONES, ONE IRON, AND THREE HIGHLY HYDROPHOBIC POLYPEPTIDE CHAINS (DESIGNATED L, M, AND H). SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN.
                                                                                                                                                                                                          SUBUNIT: REACTION CENTER IS COMPOSED OF FOUR BACTERIOCHLOROPHYLLS,
                                                                                                                                         PSBD FAMILY.
                                                                                                                                                   SIMILARITY: BELONGS TO THE REACTION CENTER L/M CHAINS / PSBA /
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ProDom; PD000551; Photo_RC; 1.
PROSITE; PS00244; REACTION_CENTER;
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15-JUL-1998 (Rel. 36, Last sequence update)
15-JUL-1998 (Rel. 36, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation update)
NADPH:ADRENODOXIN OXIDOREDUCTASE, MITOCHONDRIAL PRECURSOR
NADPH:ADRENODOXIN REDUCTASE) (AR) (FERREDOXIN-NADP(+)
(BC 1.18.1.2) (ADRENODOXIN REDUCTASE)
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BIOCHIM. BIOPHYS. AS THE FIRST ELECTRON TRANSFER PROTEIN IN ALL THE CHONCTION: SERVES AS THE FIRST ELECTRON TRANSFER PROTEIN IN ALL THE PASO SYSTEMS. INCLUDING CHOLESTEROL SIDE CHAIN MITOCHONDRIAL PASO SYSTEMS. INCLUDING CHOLESTEROLD 11-BETA CLEAVAGE IN ALL STEROIDOGENIC TISSUES, STEROID 11-BETA HYDROXYLATION IN THE ADRENAL CORTEX, 25-OH-VITAMIN D3-24 HYDROXYLATION IN THE KIDNEY, AND STEROL C-27 HYDROXYLATION IN THE KIDNEY, AND STEROL C-27 HYDROXYLATION IN THE
                                                                                                                                                                                                                                        Sagara Y., Watanabe Y., Kodama H., Aramaki H.; "cDNA cloning, overproduction and characterization of rat adrenodoxin
                                                                                                                                                                                                                                                                    STRAIN=WISTAR; TISSUE=Adrenal glan
MEDLINE=99454627; PubMed=10525147;
                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A., AND SEQUENCE OF 34-54.
                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Rodentia; Sciurognathi; Muridae; Murinae; Rat
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                                                                                                                                                                                                                Submitted (APR-2001) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                              STRAIN-CDC 1551 / Oshkosh; Fleischmann R.D., Alland D., Eisen J.A., Carpenter L., White O., Peterson J., DeBoy R., Dodson R., Gwinn M.L., Haft D., Hickey E., Kolonay J.F., Nelson W.C., Umayam L.A., Ermolaeva M.D., Salzberg S.L., Delcher A., Utterback T., Weldman J., Khouri H., Gill J., Mikula A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C., Harris D., Gordon S.V., Eiglmeier K., Gas S., Barry C.B. III, Tekaia F., Badcock K., Basham D., Brown D., Chillingworth T., Connor R., Hornsby T., Jagels K., Feltwell T., Gentles S., Hanlin N., Holroyd S., Oliver S., Osborne J., Quail M.A., Rajandream M.A., Rogers J., Stutter S., Seeger K., Skelton S., Squares S., Squares R., Steiphering the biology of Mycobacterium tuberculosis from the complete genome sequence."
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20-AUG-2001 (Rel. 40, Last annotation update)
PUTATIVE OXIDOREDUCTASE RV0945 (EC 1.-.-).
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PRINTS; PR00419; ADXRDTASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; D63761; BAA23759.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -NOV-1997 (Rel. 35, Created)
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                                                                                                                                                                                                             (SDR) FAMILY.
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                                                                                                                                                                                                                                                                                      genome comparison of Mycobacterium tuberculosis clinical and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            494 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
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58.3%;
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Pred. No. 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MITOCHONDRION
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ADRO_HUMAN
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Best Local
This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Enthe European Bioinformatics Institute. There are no restr
                                                                                                                                                                                                                                                                                                                                        "Cloning and sequence of the human adrenodoxin reductase gene.";
Proc. Natl. Acad. Sci. U.S.A. 87:8516-8520(1990).

-!- FUNCTION: SERVES AS THE FIRST ELECTRON TRANSFER PROTEIN IN ALL THE MITOCHONDRIAL P450 SYSTEMS. INCLUDING CHOLESTEROL SIDE CHAIN CLEAVAGE IN ALL STEROIDOGENIC TISSUES, STEROID 11-BETA HYDROXYLATION IN THE ADRENAL CORTEX, 25-OH-VITAMIN D3-24 HYDROXYLATION IN THE KIDNEY, AND STEROL C-27 HYDROXYLATION IN THE
                                                                                                                                                                                                                                                                   -!- COFACTOR: FAD.
                                                                                                                                                                                                                                                                                              -1- CATALYTIC ACTIVITY: REDUCED ADRENODOXIN + NADP(+) = OXIDIZED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADRO_HUMAN STANDARD; PRT; 491 AA.
P22570; 013716;
01-AUG-1991 (Rel. 19, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation update)
NADPH:ADRENODOXIN OXIDOREDUCTASE, MITOCHONDRIAL PRECURSOR
(EC 1.18.1.2) (ADRENODOXIN REDUCTASE) (AR) (FERREDOXIN-NADP(+)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
MEDLINE=91046028; PubMed=2236061;
Lin D., Shi Y., Miller W.L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chromosome 17cen-->q25 are expressed in steroidog Proc. Natl. Acad. Sci. U.S.A. 85:7104-7108(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sollsh S.B., Picado-Leonard J., Morel Y., Kuhn R.W., Mohandas T.K., Hanukoglu I., Miller W.L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=89017146; PubMed=2845396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Human adrenodoxin reductase: two mRNAs encoded by a single gene on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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                                                                                                                    PATHWAY: CHOLESTEROL SIDE-CHAIN-CLEAVAGE SYSTEM.
SUBCELLULAR LOCATION: MITOCHONDRIAL MATRIX.
ALTERNATIVE PRODUCTS: 2 ISOPORMS; A SHORT FORM (SHOWN HERE) AND A
LONG FORM; ARE PRODUCED BY ALTERNATIVE SPLICING. THE LONG FORM
REPRESENTS 10-20% OF ALL ADRENODOXIN REDUCTASE MRNA. AND SEEMS TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        les 7; Conserv
                                                                                                                 BE INACTIVE.
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Pred. No. 5
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                       a collaboration
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EMBL; J03826; AAB59497.1; -.
EMBL; M8509; AAA51668.1; -.
EMBL; M58508; AAA51668.1; JOINED.
EMBL; M58509; AAA51669.1; -.
EMBL; M58509; AAA51669.1; JOINED.
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P51524;
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01-OCT-1996 (Rel. 34, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PIG
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use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to licenseeisb-sib.ch).
                                                                                                                                                                                                                                                                                                                              Strukelj B., Pungercar J., Kopitar G., Renko M., Lenarcic B., Berbic S., Turk V.; "Molecular cloning and identification of a novel porcine
                                                                                                                                                                                                                                                                                                   cathelin-like antibacterial peptide precursor.";
Biol. Chem. Hoppe-Seyler 376:507-510(1995).
                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Bone marrow;
MEDLINE=96042752; PubMed=7576250;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sus scrofa (Pig).
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                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalia; Eutheria; Cetartiodactyla; Suina;
                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                              MEDLINE-95212585; PubMed=7698355;
                                                                                                                                                                                                                                                                             SEQUENCE OF 131-209.
                                                                                                                              "Prophenin-1, an exceptionally proline-rich antimicrobial peptide from porcine leukocytes.";
FEBS Lett. 362:65-69(1995).
1- FUNCTION: EXERTS ANTIMICROBIAL ACTIVITY. IT IS MORE EFFECTIVE AGAINST GRAM NEGATIVE BACTERIA THAN GRAM POSITIVE BACTERIA.
1- SIMILARITY: BELONGS TO THE CATHELICIDIN FAMILY.
                                                                                                                                                                                                                              Harwig S.S.L., Kokryakov V.N., Swiderek K.M., Aleshina G.M.,
                                                                                                                                                                                                                                                               TISSUE-Blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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A40487; A40487.
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7; Conserv
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n transport; Oxidoreductase; Flavoprotein; NADP; FAD;
ndrion; Transit peptide; Alternative splicing; Polymorion;
                                                                                                                                                                                                                     Lehrer R.I.
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Cetartiodactyla; Suina; Suidae;
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Pred. No. 9.6;
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E -> EALLLCQ (IN LONG ISOFORM)
O -> R.
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85865BAA2276D2B2 CRC64;
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REPEAT
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p51525;
01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
PROPHENIN-2 PRECURSOR (PF-2) (PR-2) (C12) (PROPHENIN-1 LIKE).
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Gubensek F., Turk V.;
"Molecular cloning of a putative homolog of proline/arginine-rich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota, Metázoa, Chordata, Craniata, Vertebrata,
Mammalia, Eutheria, Cetartiodactyla, Suina, Suidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sus scrofa
                                         Zhao C., Ganz T., Lehrer R.I.;
Zhao C., Ganz T., Lehrer R.I.;
"Structures of genes for two cathelin-associated antimicrobial
"Structures of genes for two cathelin-associated antimicrobial
peptides: prophenin-2 and PR-39.";
FEBS. Lett. 376:130-134(1995).
FEBS. Lett. 376:130-134(1995).
TEBS. Lett. 376:130-134(1995).
TEBS. Lett. 376:130-134 (1995).
TE
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                                                                                                                                                                                                                                                                                                       antibacterial peptides from porcine bone marrow.
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MEDLINE=94085623; PubMed=8262247;
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This SWISS-PROT entry is copyright. It is produced through a collaboration
                                                                                                                                                                                        MEDLINE=96105365; pubMed=7498526;
                                                                                                                                                                                                                         TISSUE-Liver
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                                                                                                                                                                                                                                                                                       FEBS Lett. 336:284-288(1993).
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AMIDATION (G-210 PROVIDE AMIDE GROUP)
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the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.

MEDLINE-90223989; PubMed=2158181;
Querat G., Audoly G., Sonigo P., Vigne R.;

"Nucleotide sequence analysis of SA-OMVV, a visna-related ovine
lentivirus: phylogenetic history of lentiviruses.";

Virology 175:434-447(1990).
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Pfam; PRO0666; Cathelicidin; 1.
ProDom; PD001838; Cathelicidin; 1.
PROSITE; PS00946; CATHELICIDINS 1; 1.
PROSITE; PS00947; CATHELICIDINS 2; 1.
Antibiotic; Repeat; Amidation; Signal.
Antibiotic; Repeat; Amidation; POTENTIAL.
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Viruses; Retroid viruses; Retroviridae; Lentivirus
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                                                                                                                                                          SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation -
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an email to license@isb-sib.ch).
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AMIDATION (G-226 PROVIDE AMIDE GROUP)
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Best Local
SEQUENCE FROM N.A., AND SEQUENCE OF 81-95.
MEDLINE-92015464; PubMed-1656067;
Knowles D.P. Jr., Cheevers W.P., McGuire T.C., Brassfield A.L.,
Harwood W.G., Stem T.A.;
"Structure and genetic variability of envelope glycoproteins of two
antigenic variants of caprine arthritis-encephalitis lentivirus.";
J. Virol. 65:5744-5750(1991).
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EMBL; M31646; AAA66817.1; -.
PIR; G46335; G46335.
HIV; M34193; ENV$OMVV$ACG.
InterPro; IPR000328; Env_GP41.
Pfam; PF00517; GP41; 1.
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20-AUG-2001 (Rel. 40, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation update)
ENV POLYPROTEIN PRECURSOR (COAT POLYPROTEIN) [CONTAINS: SURFACE
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                                                                                                                        ENV_CAEVC STANDARD; PRT; 966 AA.
P31626;
01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation update)
ENV POLYPROTEIN PRECURSOR (COAT POLYPROTEIN)
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Saltarelli M., Querat G., Konings D.A., Vigne R., Clements J.E. "Nucleotide sequence and transcriptional analysis of molecular of CAEV which generate infectious virus."; Virology 179:347-364(1990).
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Knowles D.P. Jr., Cheevers W.P., McGuire T.C., Brassfield A.L.,
Harwood W.G., Stem T.A.;
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pfam; PF00517; GP41; 1.
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60.0%;
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D-> P (IN REF. 2).
E-> Q (IN REF. 2).
E-> Q (IN REF. 2).
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                     Score 47; DB pred. No. 36;
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N-LINKED (GLCNAC. . .) (P
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V; 2ADEB6A92EC1D69B CRC64;
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MEDLINE-85254938; PubMed-2410140;
MEDLINE-85254938; PubMed-2410140;
Sonigo P., Alizon M., Staskus K.,
Sonigo P., Alizon M., Hasse A.,
                                                                                                                                                                                                                                        Pfam;
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MEDITINE=88062965; PubMed=2824836;
Braun M.J., Clements J.E., Gonda M.A.;
"The visna virus genome: evidence for a hypervariable site gene and sequence homology among lentivirus envelope protei J. Virol. 61:4046-4054(1987).
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p03379;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last seq)
15-JUL-1999 (Rel. 38, Last anno
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Nucleotide sequence of the visna lentivirus: relationship to the
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; M51543;
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M10608; ENV$VLVCG.
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PF00517; GP41; 1.
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CAA01216.1; -.
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POLYPROTEIN).
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Wain-Hobson S
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Andresson O.S., Elser J.E., Tobin G.J., Greenwood J.D., Gonda M.A. Georgsson G., Andresdottir V., Benediktsdottir E., Carlsdottir H.M. Maentylae E.O., Rafnar B., Palsson P.A., Casey J.W., Petursson G.; "Nucleotide sequence and biological properties of a pathogenic proviral molecular clone of neurovirulent visna virus."; Virology 193:89-105(1993).
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01-JUN-1994 (Rel. 29, Last sequence update)
20-AUG-2001 (Rel. 40, Last annotation update)
ENV POLYPROTEIN PRECURSOR (COAT POLYPROTEIN).
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L06906; AAA48362.1;
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ART PROTEIN; POLYPROTEIN; Transmembrane.

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LEADER PEPTIDE.
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EXTERIOR MEMBRANE GLYCOPROTEIN.
852
PANNEMENBRANE GLYCOPROTEIN.
852
N-LINKED (GLCNAC. . .) (POTEIN 161
N-LINKED (GLCNAC. . .) (POTEIN 162
N-LINKED (GLCNAC. . .) (POTEIN 163
N-LINKED (GLCNAC. . .) (POTEIN 164
N-LINKED (GLCNAC. . .)
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viruses; Retroviridae; Lentivirus.
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K -> AQ (IN REF. 2).

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Pred. No.
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TRANSMEMBRANE GLYCOPROTEIN.
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